

345773 Seq.ST25.txt  
SEQUENCE LISTING

10/546139

JC20 Rec'd PCT/PTO 17 AUG 2005

&lt;110&gt; Metabolic Explorer

&lt;120&gt; Procédé de préparation de microorganismes évolués permettant la création ou la modification de voies métaboliques

&lt;130&gt; D20701/ 345773

&lt;150&gt; FR 0301924

&lt;151&gt; 2003-02-18

&lt;150&gt; FR 0305768

&lt;151&gt; 2003-05-14

&lt;150&gt; FR 0305769

&lt;151&gt; 2003-05-14

&lt;150&gt; FR 0313054

&lt;151&gt; 2003-11-06

&lt;160&gt; 42

&lt;170&gt; PatentIn version 3.1

&lt;210&gt; 1

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial sequence

&lt;220&gt;

&lt;223&gt; DmetER

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## 345773 Seq.ST25.txt

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<212> DNA

<213> Artificial Sequence

<220>\_\_\_\_\_

<223> DmetEF

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<212> DNA

<213> Artificial Sequence

<220>

<223> MetER

<400> 3

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<210> 4

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> MetEF

<400> 4

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<210> 5

<211> 1161

<212> DNA

<213> Escherichia coli

<400> 5

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&lt;210&gt; 6

&lt;211&gt; 10

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 6

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1				5				10	

&lt;210&gt; 7

&lt;211&gt; 1161

&lt;212&gt; DNA

&lt;213&gt; Escherichia coli

&lt;400&gt; 7

## 345773 Seq.ST25.txt

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ctggcagaac	tggaagggtgg	tgctggtgca	gtacttacta	ataccggcat	gtccgcgatt	240
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aacggtcact	cagacgtagt	ggccggcgt	gtgattgcta	aagacccgga	cgttgtcact	660
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cggctgcaa	acaagggtt	a				1161

&lt;210&gt; 8

&lt;211&gt; 5

&lt;212&gt; PRT

&lt;213&gt; Escherichia coli

&lt;400&gt; 8

Met Glu Thr Thr His  
1 5

&lt;210&gt; 9

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

## 345773 Seq.ST25.txt

&lt;223&gt; MetJR

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&lt;210&gt; 10

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DmetJBF

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tgccagccgg aagccatttt caggcaccag agtaaacatt 100

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&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; MetCR

<400> 11  
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&lt;210&gt; 12

&lt;211&gt; 32

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; MetCF

<400> 12  
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&lt;210&gt; 13

&lt;211&gt; 100

&lt;212&gt; DNA

## 345773 Seq.ST25.txt

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DmetCR

&lt;400&gt; 13

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&lt;210&gt; 14

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DmetCF

&lt;400&gt; 14

cggacaaaaaa gcttgataact caactggtga atgcaggacg cagcaaaaaa tacactctcg 60  
gcgcggtaaa tagcgtgatt tgtaggctgg agctgcttcg 100

&lt;210&gt; 15

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DcysKR

&lt;400&gt; 15

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cccgatgatg gtagaataac catatgaata tcctccttag 100

&lt;210&gt; 16

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DcysKF

## 345773 Seq.ST25.txt

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<210> 17

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<212> DNA

<213> Artificial Sequence

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<223> DcysMR

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<211> 100

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<223> DcysMF

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<212> DNA

<213> Artificial Sequence

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<223> cysKR

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<223> cysKR

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30

<210> 21

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<212> DNA

<213> Artificial Sequence

<220>

<223> cysMR

<400> 21

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30

<210> 22

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> cysMF

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<212> DNA

<213> Artificial Sequence

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<223> MetYR

<400> 23

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gacctccaaa tgcc

345773 Seq.ST25.txt

74

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&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; MetYF

&lt;400&gt; 24

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30

&lt;210&gt; 25

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DudhAR

&lt;400&gt; 25

cccagaatct cttttgtttcccgatggAACaaaattttca gcgtgcccac gttcatgccc

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acgatttgtcgcgatgcccacatgtggctggagctgcttcg

100

&lt;210&gt; 26

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DudhAF

&lt;400&gt; 26

ggtgcgcgcg tcgcagttat cgagcgttat caaaatgttg gcggcggttg cacccactgg

60

ggcaccatcc cgtcgaaagc catatgaata tcctccttag

100

&lt;210&gt; 27

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

## 345773 Seq.ST25.txt

&lt;220&gt;

&lt;223&gt; UdhAR

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30

&lt;210&gt; 28

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&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; UdhAF

&lt;400&gt; 28

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30

&lt;210&gt; 29

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DpgiR

&lt;400&gt; 29

gcgcacacgct ttatagcggt taatcagacc attggtcgag ctatcgtggc tgctgattc

60

tttatcatct ttcagctctg catatgaata tcctccttag

100

&lt;210&gt; 30

&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; DpgiF

&lt;400&gt; 30

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## 345773 Seq.ST25.txt

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<223> PgIR

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<211> 100

<212> DNA

<213> Artificial Sequence

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<223> DpfkAR

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<212> DNA

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cgttctgcgc tgacagaagg tgtaggctgg agctgcttcg 100

&lt;210&gt; 35

&lt;211&gt; 27

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PfkAR

&lt;400&gt; 35

ccctacgccc cacttgttca tcgcccc 27

&lt;210&gt; 36

&lt;211&gt; 27

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PfkAF

&lt;400&gt; 36

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&lt;211&gt; 100

&lt;212&gt; DNA

&lt;213&gt; Artificial sequence

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&lt;223&gt; DpfkBR

&lt;400&gt; 37

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&lt;211&gt; 99

## 345773 Seq.ST25.txt

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&lt;220&gt;

&lt;223&gt; DpfkBF

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&lt;210&gt; 39

&lt;211&gt; 26

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PfkB

&lt;400&gt; 39

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&lt;210&gt; 40

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; PfkB

&lt;400&gt; 40

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&lt;210&gt; 41

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; YveDR

&lt;400&gt; 41

345773 Seq.ST25.txt

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29

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&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; YueDF

&lt;400&gt; 42

acgttcatga gatacgttat cataacagga ac

32